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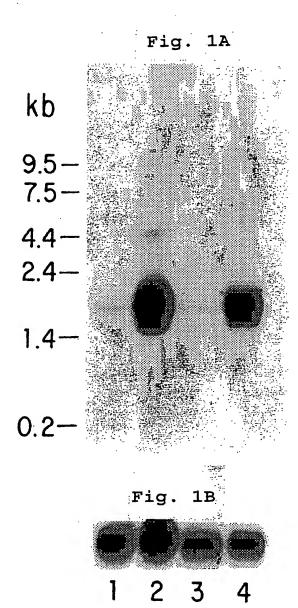
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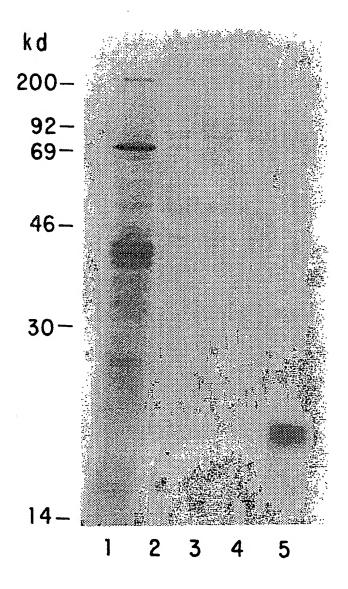


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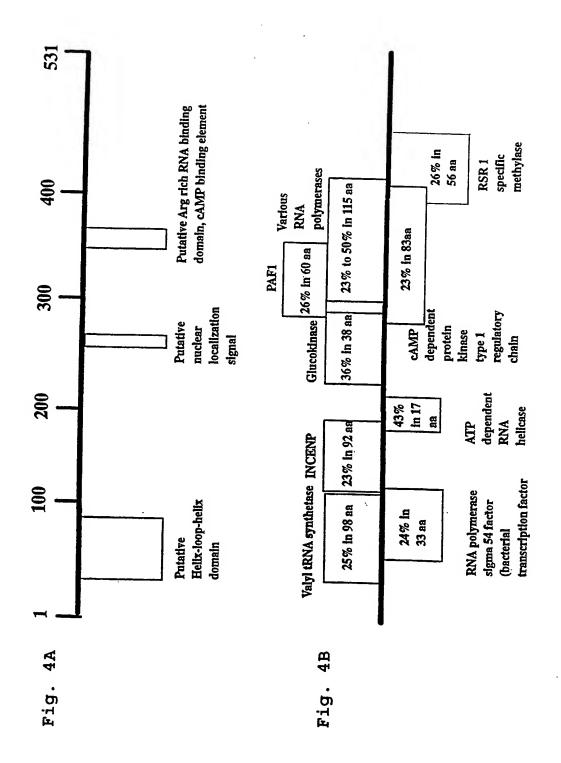
Figure 2

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61	TTCCTCGGATCAGGCGTCCCCTGAAGTCGGCACGCCCCTCTGCGTCCCCCTTCGGTCCCG	
121	CIAGGACCCGICCGGGCTGCCGTCGCCTCGTCGCTATGGCGCCCACCATCCAGACCCAG	
181	GCCCAGCGGGAGGATGGCCACAGGCCCAATTCCCACCGGACTCTGCGAGAGGTCTGGA	8
	A Y N D D G H K P N S H R T T D P D C C	
241	GTGGTCTGCCGAGTCAAGTACTGCAATAGCCTCCCTGATATCCCCTTCGACCCCAAGTTC	28
301	ATCACCTACCCCTTCGACCAGAACAGGTTCGTCCAGTACAAAGCCACTTCCTTGGAGAAA	48
361	TIPPUQNRFVOYKATSISE	68
201	CAGCACAAACATGACCTCCTGACTGAGCCAGACCTGGGGGTCACCATCGATCTCATCAAT Q H K H D L L T E P D L G V T L D L C V T L D L D L C V T L D L D L C V T L D L D L C V T L D L D L C V T L D L D L C V T L D L D L C V T L D L D L C V T L D L D L C V T L D L D L D L D L D L D L D L D L D L	
420	CCTGACACCTACCGCATCGACCCCAATGTTCTTCTAGATCCAGCTGATGAGAAACTTTTG	88
481	F D I I K I D P N V L L D P D D F K I I	108
	GAAGAGGAGATTCAGGCCCCACCAGCTCCAAGAGATCCCAGCAGCAGCAGGCGAAGGTGGTG E E I Q A P T S S K R S Q Q H A K V V	
541	CCATGGATGCGAAAGACAGAGTACATCTCCACTGAGTTCAACCGTTATGGCATCTCCAAT	128
CO1		148
601	GAGAAGCCTGAGGTCAAGATTGGGGTTTCTGTGAAGCAGCAGTTTACCGAGGAAGAAATA	240
661		168
	TACAAAGACAGGGATAGCCAGATCACAGCCATTGAGAAGACTTTTGAGGATGCCCAGAAA Y K D R D S Q I T A I E K T F E D A O K	
721	TCAATCTCACAGCATTACAGCAAACCCCGAGTCACACCGGTGGAGGTCATGCCTGTCTTC	188
701	SISUHISKPRVTPVFVMPVF	208
781	CCAGACTTTAAGATGTGGATCAATCCATGTGCTCAGGTGATCTTTGACTCAGACCCAGCC P D F K M W I N P C A C A C A C A C A C A C A C A C A C	
841		228
	CCCAAGGACACGAGTGGTGCAGCTGCGTTGGAGATGATGTCTCAGGCCATGATTAGGGGC P K D T S G A A A L E M M S O A M I R G	
901	ATGATGATGAGGAAGGGAACCAGTTTGTGGCCTATTTCCTGCCTG	248
0.61	M M D E E G N Q F V A Y F L P V F F T T	268
961	AAGAAACGAAAGCGGGACCAGGAGGAGGAGGAGATGACTATGCACCAGATGATGTGTATGAC	-50
1021	K K R K R D Q E E E M D Y A P D D V Y D	288
	TACAAAATTGCTCGGGAGTACAACTGGAACGTGAAGAACAAAGCTAGCAAGGGCTATGAG Y K I A R E Y N W N V K N K A S K G Y F	•
1081	GAAAACTACTTCATCTTCCGAGAGGGTGACGGGGTTTACTACAATGAGTTGGAAACC	308
	ENIFILF REGDGV Y N F T F m	328
1141	AGGGTCCGCCTTAGTAAGCGCCGGGCCAAGGCTGGGGTTCAGTCAG	
1201	R V R L S K R R A K A G V Q S G T N A L	348
	CTTGTGGTCAAACATCGGGACATGAATGAGAAGGAACTGGAAGCTCAGGAGGCACGGAAG L V V K H R D M N E K E L E A O E A R K	
1261	GCCCAGCTAGAAAACCACGAACCGGAGGAGAGAGAGAGAG	368
	AULENHEPEEEEEEEMETT	388
1321	AAAGAAGCTGGGGGCTCAGATGAGGAGCAGCAGGAGAAGGGCAGCAGCAGTGAGAAGGAGGGC	500
1381	REAGGS DEE OEKGSSSEKES	408
1301	AGTGAAGATGAGCACTCGGGCAGCGAGGAGGTGACAGGGACGAGGCC S E D E H S G S E S E R E E G D R D E A	
1441	AGTGACAAGAGTGGCAGTGAGGACGAGGACGAGGATGAGGCCCGGGCTGCCCGT	428
	S D A S G S G E D E S S E D E A D A A D	448
1501	GACAAAGAGGGGGTTTTTGGCAGTGATGCTGATCTGAGGACGATGCCGACTCTCATCATCATC	440
1561		468
r301	GAGGACAGAGGACAGGCCCAAGGTGGCAGTGACAATGATTCAGACAGCGGCAGCAATGGG	
1521	E D R G Q A Q G G S D N D S D S G S N G GGTGGCCAGCGGAGCCACAGCCGCAGCCCAGTCCCTTCCCCAGTGGCAGCGAG	488
	G G Q R S R S H S R S A S P F P S G S E	500
1681	CACTCGGCCCAGGAGGATGGCAGTGAAGCTGCAGCTTCTGATTCCAGTGAAGCTCATACT	508
	H S A Q E D G S E A A A S D S S E A D S	528
1741	GACAGTGACTGAGTCCCAGGGCATTCAGGGCTGGTTCAGACACCATTATTGTGAGCAGCA	520
1801	U S U *	531
1861	AAGCACTTTTCTAGTGGTCTGTTTGTGAGCCTTTCACTTGTTTGT	
921	CTTTGCTGTTAATAAAGTCAACTTCTCTTTAAAAAAAAAA	
	CHRCTITUTE CHEET (BILL E DE)	

Figure 3



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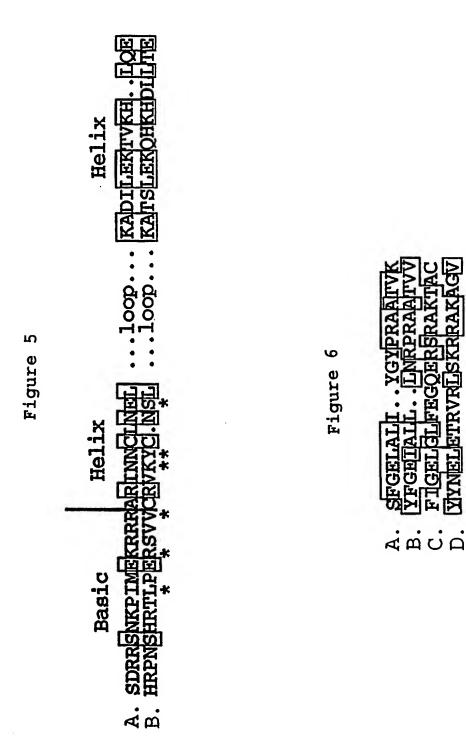
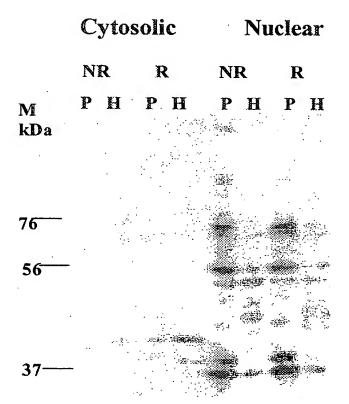
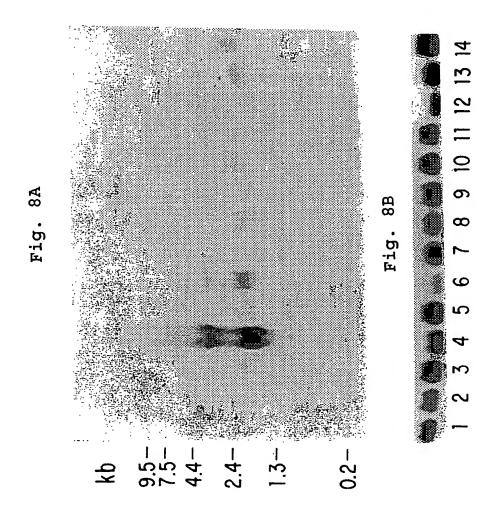


Figure 7





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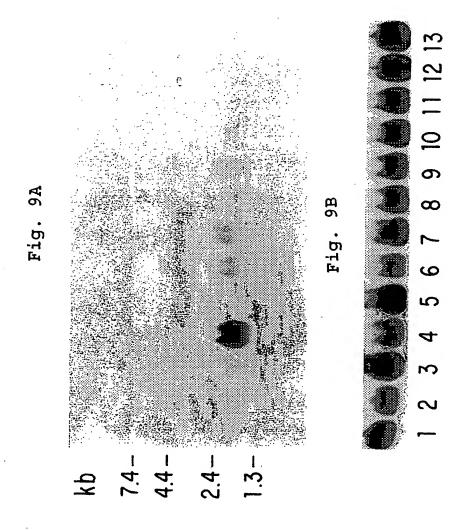


Figure 10

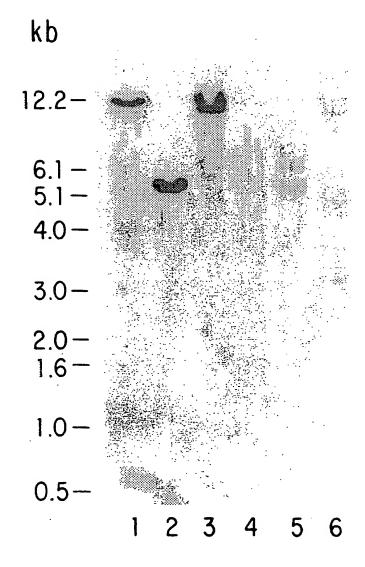
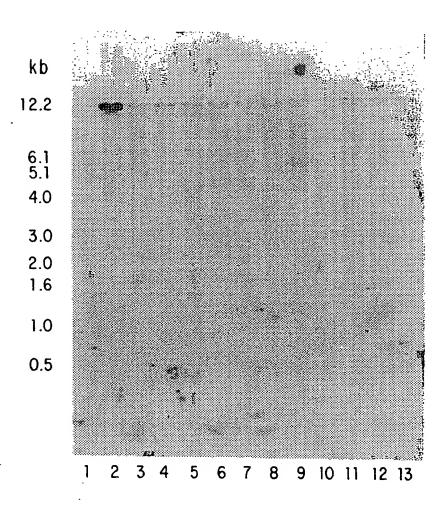


Figure 11



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Figure 12

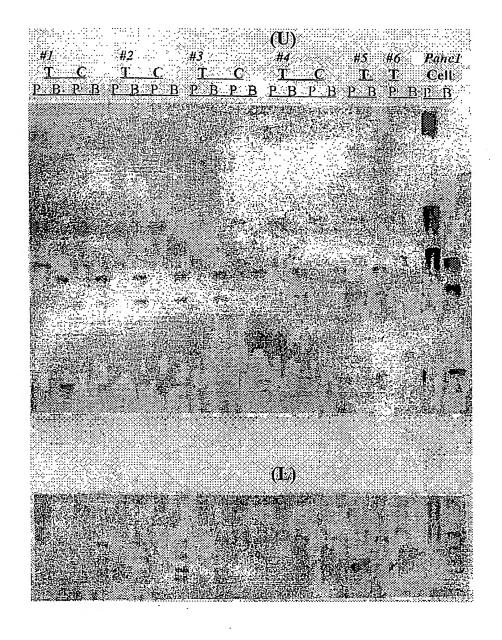
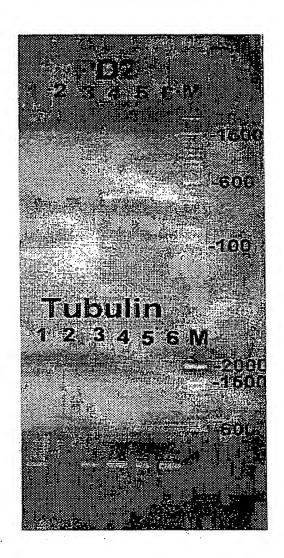


Figure 13



WO 99/50408 PCT/US99/06633

Figure 14

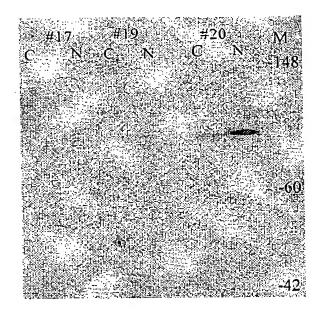


Figure 15

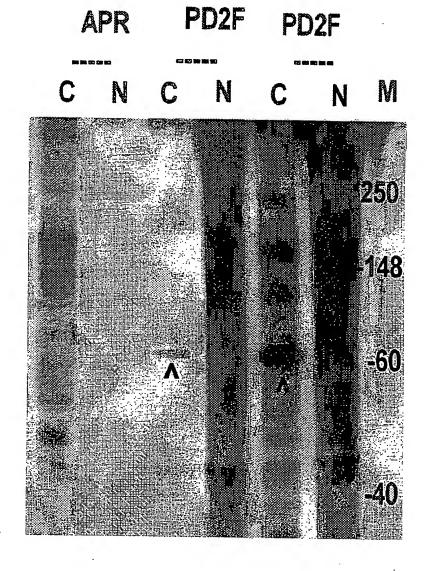


Figure 16



Figure 17

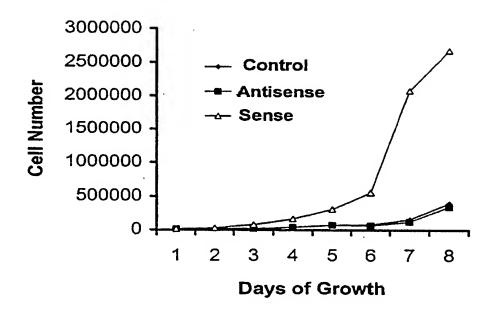


Figure 18

